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Amendment

Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims

1-7. (Canceled)

8. (Currently Amended) An expression vector comprising a plasmid selected from the group consisting of: p1-5'IL18/d1EGFP-N1 (SEO ID NO: \pm 3), p2-5'IL1 β /d1EGFP-N1 (SEQ ID NO: \pm 4), p3-5' IL1 β /d1EGFP-N1 (SEQ ID NO: $\frac{35}{5}$), p4-5'IL1 β /d1EGFP-N1 (SEQ ID NO: $\frac{46}{5}$), p1-5'3' IL1β/d1EGFP-N1 (SEQ ID NO:57), p2-5'3'IL1β/d1EGFP-N1 (SEQ ID NO:68), p3-5'3'IL1β/d1EGFP-N1 (SEQ ID NO:79), p4-5'3'IL1β/d1EGFP-N1 (SEQ ID NO:810), p1-5'IL2/EGFP-1 (SEQ ID NO:911), p1-5'IL2/d2EGFP-1 (SEQ ID NO: 1012), p1-5'3'IL2/d2EGFP-1 (SEQ ID NO: $\frac{11}{13}$), p1-3'TNF α /d1EGFP-N1 (SEQ ID NO: $\frac{12}{14}$ 14), p2-3'TNF α /EGFP-F (SEQ ID NO: ± 315), p3-3'TNF α /EGFP-F (SEQ ID NO: ± 416), p1-5'TNFα/dlegfP-N1 (SEQ ID NO: 1517), p1-5'3'TNFα/dlegfP-N1 (SEO ID NO: 1618), p1-3'IL4/d1EGFP-N1 (SEQ ID NO: 1719), p2-3'IL4/EGFP-F (SEQ ID NO: 1820), p3-3'IL4/EGFP-F (SEQ ID NO: 1921), p4-3'IL4/CA-EGFP (SEQ ID NO: 2022), p5-3'IL4/d1EGFP-N1 (SEQ ID NO: 2123), p1-5'IL4/EGFP-1 (SEQ ID NO:2224), p1-5'IL4/d1EGFP-N1 (SEQ ID NO: 2326), p2-5'IL4/EGFP-1 (SEQ ID NO: 2425), p2-5'IL4/d1EGFP-N1 (SEQ ID NO: 2527), p1-5'3'IL4/EGFP-1 (SEQ ID NO: 2630), p1-5'3'IL4/d1EGFP-N1 (SEQ ID NO:2728), p2-5'3'IL4/d1EGFP-N1 (SEQ ID NO:2829), p1-5'INFY/EGFP-1 (SEQ ID NO:2931), p1-5'INFY/d2EGFP-1 (SEQ ID NO: 3032), p1-5'3'INFY/d2EGFP-1 (SEQ ID NO: 3133), p1-5'IL10/EGFP-1 (SEQ ID NO:3237), p1-5'3'IL10/EGFP-1 (SEO ID $NO:\frac{33}{39}$), p2-5'IL10/d2EGFP-1 (SEQ ID $NO:\frac{34}{38}$), and p2-5'3'IL10/d2EGFP-1 (SEQ ID NO:3540).

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45. (Currently Amended) The expression vector of claim 8, wherein the plasmid is selected from the group consisting of: p4-5'IL1 β /d1EGFP-N1 (SEQ ID NO:46), p1-5'IL2/d2EGFP-1 (SEQ ID NO:4012), p1-5'3'TNF α /d1EGFP-N1 (SEQ ID NO:4618), p2-5'IL4/d1EGFP-N1 (SEQ ID NO:2527), p1-5'INF γ /d2EGFP-1 (SEQ ID NO:3032), and p2-5'IL10/d2EGFP-1 (SEQ ID NO:3438).

- 46. (Currently Amended) A single-celled host transformed or transfected with [[a]] the expression vector according to claim 8.
- 47. (Previously Presented) The single-celled host according to claim 46, characterised in that it is selected from the group encompassing bacteria, yeast, mammalian cells, plant cells, insect cells, as well as eukaryotic cell lines.
- 48. (Previously Presented) The single-celled host according to claim 47, characterised in that it is an immortal mammalian cell line.
- 49. (Previously Presented) The single-celled host according to claim 47, characterised in that it is a cell line selected from the group consisting of T cell leukemia cells, thymoma, mast cells, macrophage-monocytes, fibroblasts and keratinocytes.
- 50. (Previously Presented) The single-celled host according to claim 47, characterised in that it is a cell line selected from the group consisting of: EL4, BW5147.3, C57.1, J774A.1, 3T3 L1, MC/9 and HEL-30.

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51. (Previously Presented) The single-celled host according to claim 47, characterised in that it is a cell line selected from the group consisting of: C/pl-5'3'TNF α -dEGFP/2 (deposited in ECACC, Accession No. 3091202), EL/pl-5'IL2-dEGFP/6 (deposited in ECACC, Accession No. 3091204), EL/p2-5'IL4-dEGFP/2 (deposited in ECACC, Accession No. 3091205), EL/pl-5'IFN γ -dEGFP/3 (deposited in ECACC, Accession No. 3091206), EL/p2-5'IL10-dEGFP/5 (deposited in ECACC, Accession No. 3091207), and J/p4-5'IL1 β -dEGFP/4 (deposited in ECACC, Accession No. 3091208).

- 52. (Previously Presented) A method of obtaining characteristics of a tested substance, characterised in that
- a) the tested substance is put into contact with the single-celled host according to claim 46,
- b) a change in the level of expression of a green fluorescent protein caused by the tested substance is determined,
- c) the change in the level of expression described in (b) is accepted as a characteristic of the tested substance.
- 53. (Previously Presented) A collection of cell lines comprising the single celled host of claim 51 and a positive control cell line which constitutively expresses a green fluorescent protein.
- 54. (Currently Amended) The collection of cell lines of claim 53, wherein the positive control cell line is a cell line which has been transformed with a plasmid selected from the group consisting of: p1-3'GAPFH3'GAPDH/d1EGFP-N1 (SEQ ID NO:3634), p2-3'GAPHD3'GAPDH/EGFP-F (SEQ ID NO:3735), p3-3'GAPDH/EGFP-F (SEQ ID NO:3836), pCA-EGFP-F (SEQ ID NO:391), and pCA-d1EGFP (SEQ ID NO:402).

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55. (Previously Presented) The collection of cell lines of claim 53, wherein the positive control cell line is a cell line selected from the group consisting of C/pCA-EGFP-F/2 (deposited in ECACC, Accession No. 3091201) and EL/pCA-dEGFP/9 (deposited in ECACC, Accession No. 3091203).

- 56. (Previously Presented) The collection of cell lines according to claim 53, wherein the positive control cell line is a bacterial cell line, a yeast cell line, a mammalian cell line, a plant cell line, or an insect cell line.
- 57. (Previously Presented) A collection of cell lines according to claim 53, characterized in that the positive control cell line is an immortal mammalian cell line.
- 58. (Previously Presented) A collection of cell lines according to claim 53, characterized in that in the positive control cell line a gene sequence encoding the green fluorescent protein is operationally bound to a regulatory sequence selected from the group consisting of: 3 'UTR GAPDH, CMV promoter/enhancer, and actin promoter.